



1600

**Does Not Comply
Corrected Diskette Needed**

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/545,199F

DATE: 07/17/2002

TIME: 14:22:28

Input Set : A:\6227NCP.txt

Output Set: N:\CRF3\07172002\I545199F.raw

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4 <110> APPLICANT: Lowery E., David
5     Fuller E., Troy
6     Kennedy J., Michael
8 <120> TITLE OF INVENTION: Anti-Bacterial Vaccine Compositions
10 <130> FILE REFERENCE: 28341/6227.NCP
12 <140> CURRENT APPLICATION NUMBER: 09/545,199F
13 <141> CURRENT FILING DATE: 2000-04-06
15 <150> PRIOR APPLICATION NUMBER: 60/153,453
16 <151> PRIOR FILING DATE: 1999-09-10
18 <150> PRIOR APPLICATION NUMBER: 60/128,689
19 <151> PRIOR FILING DATE: 1999-04-09
21 <160> NUMBER OF SEQ ID NOS: 165
23 <170> SOFTWARE: PatentIn Ver. 2.0

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ERRORED SEQUENCES

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1994 <210> SEQ ID NO: 16
1995 <211> LENGTH: 2110
1996 <212> TYPE: PRT
1997 <213> ORGANISM: Pasteurella multocida
1999 <400> SEQUENCE: 16
2000 Met Gln Pro Ala Gln Glu His Cys Gln Arg Ile Asn Asn Ile Val Asn
2001      1           5           10          15
2003 Gln Glu Asn Gly Leu Phe His Thr Leu Gly Asn Met Met Leu Glu Ala
2004      20          25          30
2006 Glu Arg Ser Val Tyr Asn Ile Gly Asp Ile Tyr Ala Ser Lys Lys Leu
2007      35          40          45
2009 Thr Val His Thr His Asn Leu Ile Asn Asp Val Arg Leu Ser Gly Asn
2010      50          55          60
2012 Val Ser Tyr Lys Pro Ile Gly Ser Ser Arg/Asp Tyr Asp Ile Ser Arg
2013      65          70          75          80
2015 Val Ala Val His Gly Trp His Asn Asn Val Tyr Lys Leu Asn Leu Asn
2016      85          90          95
2018 Leu Gln Glu Gln Asp Lys Thr Asp Ile Lys Val Val Lys Met Gly Ala
2019      100         105         110
2021 Ile Arg Ser Asp Gly Asp Phe Asp Phe Lys Gly Ile Lys Ala Thr Ser
2022      115         120         125
2024 Ser Glu Ser Lys Pro Gln Leu Ile Asn His Gly Leu Ile Asn Val Lys
2025      130         135         140
2028 Gly Thr Phe Asn Ala Glu Ala Asp Gln Val Val Asn Gln Met Lys Ala
2029 145           150           155           160
2031 Phe Asn Gln Asn Ala Leu Ala Ser Val Phe Lys Asn Pro Ala Lys Ile

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2032	165	170	175
2034	Thr Met Tyr Tyr Gln Pro Leu Thr Arg Tyr Ile Trp Thr Pro Leu Ser		
2035	180	185	190
2037	Gly Asn Ala Ser Arg Glu Phe Asn Asn Leu Glu Ser Phe Leu Asp Ala		
2038	195	200	205
2040	Leu Phe Gly Ser Thr Thr Ile Leu Lys Ser Ser Phe Tyr Ser Thr Glu		
2041	210	215	220
2043	Asn Phe Ser Ala Tyr Gln Leu Leu Ser His Ile Gln His Ser Pro Met		
2044	225	230	235
2046	Tyr Gln Lys Ala Met Ala Gln Val Phe Gly Ala Glu Trp His Ser Lys		
2047	245	250	255
2049	Ser Tyr Asp Glu Met Arg Asn Lys Trp Lys Ser Phe Lys Glu Asn Pro		
2050	260	265	270
2052	Thr Asp Phe Ile Tyr Tyr Pro Ser Glu Lys Ala Lys Ile Leu Ala Gly		
2053	275	280	285
2055	Lys Leu Glu Gly Lys Leu Thr Thr Leu Gln Asn Gly Glu Tyr Ala Glu		
2056	290	295	300
2058	Arg Gly Lys Phe Asp Glu Ser Ile Gln Ile Gly Lys His Gln Leu Ser		
2059	305	310	315
2061	320		
2062	Leu Pro Ser Val Glu Leu Lys Ala Glu Phe Ser Asp Lys Glu Arg Leu		
2064	325	330	335
2065	Glu Glu Asp Gly Val Asp Leu Ser Ser Ile Ala Glu Leu Leu Glu Met		
2067	340	345	350
2068	Pro Asn Leu Phe Ile Asp Asn Ser Ile Gln Leu Glu Lys Lys Lys Leu		
2070	355	360	365
2071	Ser Pro Ile Glu Asp Leu Asp Glu Glu Pro Arg Lys Asn Leu Asp Ile		
2073	370	375	380
2074	Glu Glu Ser His Ser Asn Ser Ser Asp Asp Val Leu Ser Met Asn Asp		
2075	385	390	395
2076	400		
2077	Asp Glu Ser Asp Thr Asp Asp Ser Lys Trp Ser Met Gly Asn Asp Glu		
2079	405	410	415
2080	Lys Glu Met Pro Asp Asp Lys Leu Gly Ile Ser Arg Asp Asp Arg Gly		
2082	420	425	430
2083	Asn Lys Pro Pro Arg Thr Asp Pro Thr Val Asp Tyr Leu Asn Pro Asp		
2085	435	440	445
2086	Glu Phe Glu Asn Gly Tyr Leu Leu Asn Glu Leu Leu Gln Glu Leu		
2089	450	455	460
2090	Gly Glu Glu Pro Leu Leu Lys Glu Gly Glu Asp His Phe Lys Arg Ser		
2092	465	470	475
2093	480		
2095	Thr Asn Leu Val Arg Leu Gly Glu Arg Asp Arg Gln Asn Arg Glu Lys		
2096	485	490	495
2098	Arg Glu Lys Glu Gly Tyr Phe Asp Leu Pro Gly Thr Leu Asp Met Lys		
2099	500	505	510
2101	Leu Gln Glu Leu Phe Glu Lys Arg Lys Gln Lys His Glu Ala Glu Gln		
2102	515	520	525
2104	Lys Ala Arg Ile Glu Lys Ala Leu Leu Gln Lys Ser Glu Gln Gln Glu		
2105	530	535	540
	545	550	555
			560

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2107 Lys Ile Ala Lys Gln Val Glu Ile Ala Lys Glu Met Gln Arg Val Glu
2108      565      570      575
2110 Glu Ile Arg Gln Arg Glu Lys Gln Leu Ala Ile Gln Leu Gln Glu Glu
2111      580      585      590
2113 Glu Lys Lys Gln Gln Glu Glu Lys His Leu Ser Glu Glu Lys Lys Gln
2114      595      600      605
2116 Ala Glu Gln Lys Gln Lys Ala Glu Glu Lys Val Ala Gln Glu Arg Leu
2117      610      615      620
2119 Asp Ile Glu Gln Gln Lys Ala Tyr Glu Glu Met Ala Lys Arg Glu Ala
2120 625      630      635      640
2122 Glu Ala Ser Lys Asn Val Leu Leu Lys Ala Ile Asp Glu Glu Arg Pro
2123      645      650      655
2125 Lys Val Glu Thr Asp Pro Leu Phe Arg Thr Lys Leu Lys Tyr Ile Asn
2126      660      665      670
2128 Gln Asp Asp Tyr Ala Gly Ala Asn Tyr Phe Phe Asn Lys Val Gly Leu
2129      675      680      685
2131 Asn Thr Lys Gly His Gln Lys Val Asn Val Leu Gly Asp Asn Tyr Phe
2132      690      695      700
2134 Asp His Gln Val Ile Thr Arg Ser Ile Glu Lys Lys Val Asp Asn His
2135 705      710      715      720
2137 Leu Asn Gln Lys Tyr Asn Leu Ser Asp Val Glu Leu Val Lys Gln Leu
2138      725      730      735
2140 Met Asp Asn Ser Thr Thr Gln Ala Gln Glu Leu Asp Leu Lys Leu Gly
2141      740      745      750
2143 Ala Ala Leu Thr Lys Glu Gln Ala Asn Leu Thr Gln Asp Ile Val
2144      755      760      765
2146 Trp Tyr Val Lys Thr Lys Val Lys Gly Lys Asp Val Phe Val Pro Lys
2147      770      775      780
2150 Val Tyr Phe Ala Ser Glu Thr Leu Val Glu Ala Gln Lys Leu Gln Gly
2151 785      790      795      800
2153 Leu Gly Thr Gly Thr Ile Arg Val Gly Glu Ala Lys Ile Lys Ala Lys
2154      805      810      815
2156 Asp Val Val Asn Thr Gly Thr Leu Ala Gly Arg Lys Leu Asn Val Glu
2157      820      825      830
2159 Ala Ser Asn Lys Ile Lys Asn Gln Gly Ser Ile Leu Ser Thr Gln Glu
2160      835      840      845
2162 Thr Arg Leu Val Gly Arg Lys Gly Ile Glu Asn Val Ser Arg Ser Phe
2163      850      855      860
2165 Ala Asn Asp Glu Leu Gly Val Thr Ala Gln Arg Ser Glu Ile Lys Thr
2166 865      870      875      880
2168 Glu-Gly His Leu His Leu Glu Thr Asp Lys Asp Ser Thr Ile Asp Val
2169      885      890      895
2171 Gln Ala Ser Asp Ile Lys Ala Lys Thr Ser Phe Val Lys Thr Gly Asp
2172      900      905      910
2174 Val Asn Leu Lys Asn Thr Tyr Asn Thr Lys His Ala Tyr Arg Glu Lys
2175      915      920      925
2177 Phe Ser Pro Ser Ala Leu Gln Val Ala Glu Leu Asp Val Ala Gly Leu
2178      930      935      940
2180 Lys Val Pro Leu Leu Gly Val Ser Ser Pro Ser Tyr Ser Glu His

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2181	945	950	955	960
2183	Thr Ser Glu Ala Thr Ser Glu Gly Ser Ile Phe Glu Val Gly His Leu			
2184	965	970	975	
2186	His Leu Ala Val Asp Arg Asp Val Asn Gln Ala Gly Ser Lys Ile Lys			
2187	980	985	990	
2189	Ala Lys Tyr Thr Thr Gly Val Val Lys Gly Asn Phe Asn Thr Glu Ala			
2190	995	1000	1005	
2192	Gly Lys Asn Ile Lys His Val Glu Lys Glu Glu Tyr Ser Ser Gln Leu			
2193	1010	1015	1020	
2195	Phe Ala Ser Ala His Ala Ser Gly Gly Thr Ser Val Arg Tyr Asp			
2196	1025	1030	1035	1040
2198	Tyr Asn Ser Gln Asp Gly Gly Asn Ala Ser Val Gly Val Pro Thr Asn			
2199	1045	1050	1055	
2201	His Thr Gly Val Gly Ala Glu Ala Gly Met Ser Phe Thr His Thr Lys			
2202	1060	1065	1070	
2204	Asp Lys Glu Thr Leu Leu Thr His Thr Asn Ser Glu Leu Gln Val Lys			
2205	1075	1080	1085	
2207	His Gly Lys Leu His Val Leu Gly Tyr Ala Asp Ile Gly Gly Val Asp			
2208	1090	1095	1100	
2211	Ile Asn Thr Lys Leu Pro Glu Asp Ala Gln Ser Lys Ala Gln Lys Glu			
2212	1105	1110	1115	1120
2214	Ile Ala Ala Ser Lys Pro Glu Lys Thr Glu Gln Ser Ala Gln Asp Val			
2215	1125	1130	1135	
2217	Ala Gln Ala Gln Ser Asn Ala Asn Lys Asp Lys Glu Asn Lys Ala Pro			
2218	1140	1145	1150	
2220	Glu Ile Lys Glu Leu Ser Glu Ala Glu Ile Ala Asp Leu Met Ser Glu			
2221	1155	1160	1165	
2223	Lys Ser Lys Ala Tyr Phe Asp Asp Phe Ala Glu Gln Ala Lys Lys Ala			
2224	1170	1175	1180	
2226	Pro Glu Asn Asn Arg Phe Glu Leu Ser Ala Lys Glu Ile Lys Ser Ser			
2227	1185	1190	1195	1200
2229	Lys Gln Lys Asp Gln Tyr Asp His Glu Ser Glu Arg Thr Thr Phe Lys			
2230	1205	1210	1215	
2233	Val Gly Pro Glu Ala Glu Ala His Ser Ala Val Ala Asp Met Val Ser			
2234	1220	1225	1230	
2236	His Leu Val Lys Glu Tyr Arg Asp Ala Gln Asn Gly Thr Lys Gln Asp			
2237	1235	1240	1245	
2239	Gly Thr Val Ala Leu Gln His Ala Ser Asp Val Leu Asn Ile Val Thr			
2240	1250	1255	1260	
2242	Gly Asp Leu Ala Gly Ser Ser Ala Lys Leu Ser Val Glu Arg Thr His			
2243	1265	1270	1275	1280
2245	Glu Thr Lys Arg Thr Thr Glu Thr Gly Asp Ile Val Thr Lys Ile Gly			
2246	1285	1290	1295	
2248	Gly Asn Val Thr Leu Ser Ala Arg Ser Gly Ser Val Asn Leu Lys Asn			
2249	1300	1305	1310	
2251	Val Gln Ser Asp Glu Gln Ala Asn Leu Thr Leu Arg Ala Lys Glu Asp			
2252	1315	1320	1325	
2254	Val Asn Val Leu Ser Gly Glu Lys Thr Arg Glu Thr Thr Glu Thr Val			
2255	1330	1335	1340	

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2257	Ser Arg Gln Lys Leu Ser His Gly Val Asn Ala Gly Cys Ser Met Met			
2258	1345	1350	1355	1360
2260	Ser Gly Ala Cys Thr Ala Gly Val Ser Thr Ser Leu Glu Gly Asn Glu			
2261	1365	1370	1375	
2263	Ser Tyr Thr Ser Glu Arg Glu Thr Ala Gln Asn Asn Ser Phe Leu Lys			
2264	1380	1385	1390	
2266	Ala Arg Asn Met Lys Val Glu Ala Gly Arg Asp Phe Asn Val Val Ser			
2267	1395	1400	1405	
2269	Ser Asn Ile Asp Ala Asp Lys Leu Asp Leu His Val Lys Gly Lys Thr			
E-->	2270 A1410	1415	1420	
2272	Asn Val Val Ser Lys Gln Asp Thr Leu Gln Lys Val Thr His Gly Val			
E-->	2273 1425	1430	1435	←1440
2275	Asp Tyr Asn Leu Ser Ala Gly Val Ala Leu Ser Ser Ala Thr Ile Ala			
E-->	2276 1445	1450	1455	
2278	Thr Pro Thr Gly Asn Val Gly Phe Gly Tyr Thr Asn Glu Thr Glu Ser			
E-->	2279 1460	1465	1470	
2281	Lys Arg Thr Val Asn Gln Gln Ala Gly Ile Lys Ala Asn Lys Ile Thr			
E-->	2282 1475	1480	1485	
2284	Gly Gln Thr His Asp Leu Asn Leu Glu Gly Gly Tyr Leu Val Ser Asn			
E-->	2285 1490	1495	1500	
2287	Asp Lys Asp Asn Gln Leu Lys Val Thr Gly Asp Val Thr Thr Lys Ala			
E-->	2288 1505	1510	1515	←1520
2290	Leu His Asp Gln His Asp Lys Asp Gly Gly Thr Phe Gly Leu Ser Val			
E-->	2291 1525	1530	1535	
2293	Gly Ile Ser Glu Arg Gly Thr Thr Ala Phe Asn Val Arg Gly Gly Arg			
E-->	2294 1540	1545	1550	
2296	Ala Glu Gln Lys His Tyr Asn Ala Thr Gln Lys Ser Thr Leu Ser Gly			
E-->	2297 1555	1560	1565	
2299	Val Asp Thr Ser Gln Ala Asn Val Ser Gly Gln Val Asn Thr Asp Leu			
E-->	2300 1570	1575	1580	
2302	Thr Lys Ala Lys Ala Val Thr Arg Asp Asp Thr Tyr Ala Ser Thr Gln			
E-->	2303 1585	1590	1595	←1600
2305	Phe Ser Phe Glu Val Ala Asp Ile Val Glu Leu Gly Gln Arg Ala Lys			
E-->	2306 1605	1610	1615	
2308	Asn Lys Leu Ser Ala Pro Asn Asn Asp Thr Asp Met Ala Ser Gly Ser			
E-->	2309 1620	1625	1630	
2311	Thr Leu Arg Ser Arg Ser Thr Thr Glu Glu Ala Asp Val Pro Thr Thr			
E-->	2312 1635	1640	1645	
2314	Arg Ser Arg Val Thr Asp Glu Ala Asp Ser Val Ser Val Lys Asn Pro			
E-->	2315 1650	1655	1660	
2317	Ile Tyr Glu Ser Ala Asp Ala Val Val Pro Thr Pro Arg Ser Arg Asn			
E-->	2318 1665	1670	1675	←1680
2320	Val Asp Ser Thr Asp Leu Val Asp Asn Pro Leu Tyr Ala Ser Ala Thr			
E-->	2321 1685	1690	1695	
2323	Thr Lys Ala Asn Ile His Asp Tyr Glu Glu Ile Pro Ala Val Tyr Ser			
E-->	2324 1700	1705	1710	
2326	Lys Val Gly Asp Asn Asn Ala Asp Leu Val Arg His Lys Thr Ala Thr			
E-->	2327 1715	1720	1725	
2329	Ser Asp Glu His Leu Tyr Ala Glu Ile Asn Glu Pro Thr Tyr Ser Arg			

amino number
misaligned

- move to the left one
space, so that the last
digit aligns with the
right margin (last
letter of amino
acid)

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E--> 2330	1730	1735	1740
2333	Val Gly Asp Lys Asn Ala Asp Met Arg Arg His Asn Ala Ala Gly Thr		
E--> 2334	1745	1750	1755
2336	Thr Asp Tyr Ala Asp Val Val Gln Ala His Thr Arg Lys Ala Asp Asp		↳1760
E--> 2337		1765	1770
2339	Pro Leu Pro Ala Leu Pro Asn Gln Gly Lys Ala Arg Thr Val Asn Asp		1775
E--> 2340		1780	1785
2342	Gly Ser Glu His Ile Tyr Thr Asp Ile Ser Asp Val Gly Thr Gln Thr		
E--> 2343		1795	1800
2345	Lys Ala Ile Asp Ser Thr Tyr Ala Thr Val Gly Met Pro Lys Ala Asn		1805
E--> 2346		1810	1815
2348	Ala Val Asn Leu Ile Gly Gln Asn Gly Leu Gly Ser Ile Tyr His Ser		1820
E--> 2349		1825	1830
2351	Pro Asp Ser Ala Tyr Lys Thr Trp Gln Leu Leu Asp Gln Phe Ala Asn		↳1840
E--> 2352		1845	1850
2354	Lys Gly Gly Asp Ala Val Phe Leu Arg Pro Ala Thr Glu Met Lys Cys		1855
E--> 2355		1860	1865
2358	Ala Gly Ala Pro Leu Lys Tyr Thr Phe Ile Val Arg Asp Tyr Leu Leu		1870
E--> 2359		1875	1880
2361	Arg Arg His Thr Leu Asp Lys Ser Arg Leu Phe Tyr Asn Ala His Asn		1885
E--> 2362		1890	1895
2364	Lys Thr Leu Phe Ser Val Pro Ile Val Asp Ala Lys Val Lys Met Leu		1900
E--> 2365		1905	1910
2367	Phe Ala Glu Lys Asn Ile Gln Val Asn Tyr Asp Arg Ser Leu Thr Ala		↳1920
E--> 2368		1925	1930
2370	Ile Asp Leu Ser Lys Arg Ile Ala Thr Phe Asn Ser Pro Glu Gly Val		1935
E--> 2371		1940	1945
2373	Val Glu Val Pro Tyr Asp Phe Ile Asn Val Val Pro Pro Met Arg Ala		1950
E--> 2374		1955	1960
2376	Pro Asp Ala Val Arg Gln Ser Ala Leu Ala Trp Gln Glu Gly Lys Trp		1965
E--> 2377		1970	1975
2379	Ala Asn Asp Gly Trp Val Glu Val Glu Lys His Thr Leu Arg His Arg		1980
E--> 2380		1985	1990
2382	Arg Tyr Ala Asn Val Phe Ala Val Gly Asp Val Ala Gly Val Pro Lys		↳2000
E--> 2383		2005	2010
2385	Gly Lys Thr Ala Ala Ser Val Lys Trp Gln Val Pro Val Ala Val Ala		2015
E--> 2386		2020	2025
2388	His Leu Leu Ala Glu Leu Glu Gly Lys Pro Cys Asp Glu Ile Tyr Asn		2030
E--> 2389		2035	2040
2391	Gly Tyr Thr Ser Cys Pro Leu Ile Thr Gln Leu Gly Lys Gly Met Leu		2045
E--> 2392		2050	2055
2394	Val Glu Phe Asp Tyr Asn Asn His Leu Thr Pro Ser Phe Pro Gly Val		2060
E--> 2395		2065	2070
2397	Ile Ala Pro Leu Glu Glu Leu Trp Ala Thr Trp Ala Ile Lys Thr Leu		↳2080
E--> 2398		2085	2090
2400	Gly Leu Lys Pro Thr Tyr Leu Gly Met Leu Arg Gly Leu Ala		2095
E--> 2401		2100	2105
			2110

VERIFICATION SUMMARY
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L:129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:1061
L:2270 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:16
M:332 Repeated in SeqNo=16
L:3841 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:337
L:3842 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:385
L:3845 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:385
L:3846 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:433
L:3849 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:433
L:3850 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:481
L:3981 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:112
L:3984 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:128
L:3987 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:144
L:5416 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:975
L:5417 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1023
L:5428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1119
L:5429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1167
L:5444 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1390
L:5452 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:1630
L:5526 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:208
L:5529 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:224
L:5535 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:256
L:5538 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:272
L:5612 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:97
L:5613 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:145
L:5628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:289
L:5629 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:337
L:5661 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:889
L:5663 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:949
L:5689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:1729
L:5691 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:1789
L:5728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:32
L:5740 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:96
L:5858 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:1806
L:6790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47 after pos.:3076
L:9359 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:72 after pos.:420
L:11243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:90 after pos.:0
L:11245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:90 after pos.:60
L:11978 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:102 after pos.:4848
L:11979 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:102 after pos.:4896
L:12306 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:103 after pos.:1616